

SEQUENCE LISTING

<110> Paul Young et al.

<120> PGRP-L Polynucleotides, Polypeptides, and Antibodies

<130> PF513P1

<150> 60/149,715

<151> 1999-08-20

<150> PCT/US00/22877

<151> 2000-08-18

<160> 18

<170> PatentIn version 3.1

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35 40 45

Arg Ala Gly Leu Leu Arg Pro Asp Tyr Ala Leu Leu Gly His Arg Gln
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Leu Val Arg Thr Asp Cys Pro Gly Asp Ala Leu Phe Asp Leu Leu Arg
65 70 75 80

Thr Trp Pro His Phe Thr Ala Val Ser Leu Arg Ser Leu His Tyr Thr
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Ala Arg Arg Pro Ser Val Tyr Thr Ser Ser Thr Arg Pro Leu Pro Pro
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Ala Cys Asn Ser Cys Ala Arg Thr Ala Ser Ala Arg Pro Pro Thr Ser
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Arg Arg His Val Tyr Ser Gly Asn Leu Gly Pro Ala Phe Ala Gly His
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1876

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Ile Gln Ala Leu Ala Glu Leu Glu Gln Lys Val Pro Val Thr Glu Ala
35 40 45

Ser Ile Thr Ala Ser Ala Trp Ile Leu Ser Ala Lys Asn Ser Ser Thr
50 55 60

His Asn Ser Leu His Gln Arg Leu Leu Leu Lys Ala Pro Ser His Asn
65 70 75 80

Thr Thr Glu Pro Asp Pro His Ser Leu Ser Pro Glu Leu Gln Ala Leu
85 90 95

Ile Ser Glu Val Ala Gln His Asp Val Gln Asn Gly Arg Glu Tyr Gly
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Val Val Leu Ala Pro Asp Gly Ser Thr Val Ala Val Lys Pro Leu Leu
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Phe Gly Leu Glu Ala Gly Leu Gln Ala His Ser Val Ala Asn Leu Pro
130 135 140

Ser Asp Cys Leu Ala Ile Pro Cys Asp Thr Gly Asp Thr Leu Ala Asn
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Ile Arg Ala Thr Trp Pro Gly Leu Met Asp Ala Phe Pro Asn Ala Ser
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Ser Pro Asp Val Gly Ala Thr Leu Pro Asn Asp Lys Ala Lys Thr Pro
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Thr Thr Val Asp Arg Leu Leu Ala Ile Thr Leu Ala Gly Asp Leu Gly
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Thr Glu Gly Cys Trp Asp Gln Leu Thr Ala Pro Arg Val Phe Thr Leu
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Leu Asp Pro Gln Ala Ser Arg Leu Thr Met Ala Phe Leu Asn Gly Ala
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His Pro Pro Leu Ser His Leu Leu Arg Glu Tyr Tyr Gly Ala Gly Val
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Asn Gly Asp Pro Val Phe Arg Ser Asn Phe Arg Arg Gln Asn Gly Ala
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Ala Leu Thr Ser Ala Pro Thr Leu Ala Gln Gln Val Trp Glu Ala Leu
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Val Leu Leu Gln Lys Leu Glu Pro Glu His Leu Gln Leu Gln Asn Ile
325 330 335

Ser Gln Glu Gln Leu Ala Gln Val Ala Thr Leu Ala Thr Lys Glu Phe
340 345 350

Thr Glu Ala Phe Leu Gly Cys Pro Ala Ile His Pro Arg Cys Arg Trp
355 360 365

Gly Ala Ala Pro Tyr Arg Gly His Pro Thr Pro Leu Arg Leu Pro Leu
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Gly Phe Leu Tyr Val His His Thr Tyr Val Pro Ala Pro Pro Cys Thr
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Thr Phe Gln Ser Cys Ala Ala Asp Met Arg Ser Met Gln Arg Phe His
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Gln Asp Val Arg Lys Trp Asp Asp Ile Gly Tyr Ser Phe Val Val Gly
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Ser Asp Gly Tyr Leu Tyr Gln Gly Arg Gly Trp His Trp Val Gly Ala
435 440 445

His Thr Arg Gly Tyr Asn Ser Arg Gly Phe Gly Val Ala Phe Val Gly
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Asn Tyr Thr Gly Ser Leu Pro Asn Glu Ala Ala Leu Asn Thr Val Arg
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Asp Ala Leu Pro Ser Cys Ala Ile Arg Glu Gly Leu Leu Arg Pro Asp
485 490 495

Tyr Lys Leu Leu Gly His Arg Gln Leu Val Leu Thr His Cys Pro Gly
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Glu Asn
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gccctaact ccgcccagtt ccgcccattc tccgccccat ggctgactaa ttttttttat 180

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<211> 31
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